### Heritable Disorders of Connective Tissue Panel

#### Panel Gene List

ACTA2, ADAMTS2, AEBP1, ALDH18A1, ATP6V0A2, ATP6V1E1, ATP7A, B3GALT6, B3GAT3, B4GALT7, BGN, CBS, CHST14, COL1A1, COL1A2, COL2A1, COL3A1, COL4A1, COL5A1, COL5A2, COL9A1, COL9A2, COL9A3, COL11A1, COL11A2, COL12A1, DSE, EFEMP2, ELN, FBLN5, FBN1, FBN2, FKBP14, FLNA, LOX, LTBP4, MAT2A, MED12, MFAP5, MYH11, MYLK, NOTCH1, PLOD1, PRDM5, PRKG1, PYCR1, RIN2, SKI, SLC2A10, SLC39A13, SMAD2, SMAD3, SMAD4, TAB2, TGFB2, TGFB3, TGFBR1, TGFBR2, TNXB, ZNF469

#### **Clinical Features:**

The heritable disorders of connective tissue (HDCT) are a group of clinically and genetically heterogeneous conditions that variably involve the cardiovascular, musculoskeletal, cutaneous, ocular, craniofacial, pulmonary, gastrointestinal and/or neurologic systems.<sup>1</sup> Overlapping features and variable expressivity may challenge clinical diagnosis.<sup>2,3,4</sup>

Cardiovascular complications that occur in some HDCTs include aortic and arterial aneurysm, dissection or rupture, as well as arterial tortuosity and stenosis, mitral valve prolapse, and patent ductus arteriosus. An increased risk for thoracic aortic aneurysm and dissection (TAAD) may be associated with other syndromic features in Marfan syndrome, Loeys-Dietz syndrome (LDS), Shprintzen-Goldberg syndrome and vascular Ehlers-Danlos syndrome (vEDS).<sup>5</sup> Aortopathy also occurs more frequently in individuals with pathogenic variants in a number of other genes, including

ACTA2, BGN, LOX, MAT2A, MFAP5, MYH11, MYLK, NOTCH1, PRKG1 and SMAD4.<sup>3,6,7,8</sup> Aneurysms, tortuosity, dissections and/or rupture of other arteries can occur in vEDS, LDS, arterial tortuosity syndrome (ATS), and other rare forms of Ehlers-Danlos syndrome.<sup>9,10,11</sup> Supravalvular aortic stenosis may be due to pathogenic variants in *ELN*.<sup>12</sup>

Musculoskeletal features in HDCTs include joint hypermobility, scoliosis and/or kyphosis, pectus excavatum/carinatum, precocious osteoarthritis and/or various forms of skeletal dysplasias. Congenital hip dislocation is a hallmark feature of arthrochalasia type EDS<sup>13</sup> but may also occur in other HDCTs. Craniosynostosis may occur in LDS<sup>10</sup> or Sphrintzen-Goldberg syndrome.<sup>14</sup> Contractures may be seen in individuals with congenital contractural arachnodactyly, musculocontractural EDS, or ATS. Stature may be tall in individuals with Marfan syndrome or homocystinuria and short in individuals with Stickler syndrome, or osteogenesis imperfecta (OI).

Ocular complications are associated with some HDCTs. High myopia with or without retinal detachment is a feature of Stickler syndrome or Marfan syndrome. Ectopia lentis occurs in Marfan syndrome and homocystinuria.<sup>15</sup> Keratoconus can occur in individuals with vEDS, brittle cornea syndrome (BCS), or ATS.<sup>9</sup> Iris floculi may be associated with pathogenic variants in *ACTA2*.<sup>16</sup> Individuals with BCS or kyphoscoliotic EDS are at risk of spontaneous perforation of the eye.<sup>13,17</sup>

The cutaneous features of HDCTs include relatively common findings, such as striae and easy bruising, or less frequently seen characteristics, such as hyperextensible or doughy skin, skin fragility and atrophic scars. Loose inelastic skin is a key finding in cutis laxa but also occurs in other conditions, such as ATS or occipital horn

syndrome (OHS).<sup>9</sup> Increased skin transparency may occur in vEDS, LDS, or other rare forms of EDS. Livedo reticularis may be associated with pathogenic variants in *ACTA2*.<sup>16</sup>

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Complications of HDCTs may also occur in other systems. Pneumothorax with or without pulmonary blebs are seen in Marfan syndrome and vEDS. Spontaneous bowel and hollow organ rupture can occur in persons with vEDS, as well.<sup>11</sup> Hearing loss is associated with Stickler syndrome, OI, BCS, and FKBP14-related EDS.<sup>17,18,19</sup> Other neurological complications of some HDCTs include hypotonia, myopathy, periventricular heterotopia with or without seizures (*FLNA*-related disorders<sup>20</sup>), developmental delays or intellectual disability (Sprintzen-Goldberg syndrome, homocystinuria) or psychiatric illness (homocystinuria).<sup>21,22</sup>

For more in-depth information on a specific HDCT, please refer to OMIM or GeneReviews for the condition of interest or to the references cited above. SECTION 3

#### **Genetics:**

Autosomal Dominant, Autosomal Recessive, X-linked

#### **Test Methods:**

Using genomic DNA extracted from the submitted specimen, the complete coding regions and splice site junctions of the genes tested are enriched using a proprietary targeted capture system developed by GeneDx for next-generation sequencing with CNV calling (NGS-CNV) (only exons 1-

31 for TNXB). The enriched targets are simultaneously sequenced with paired-end reads on an Illumina platform. Bi-directional sequence reads are assembled and aligned to reference sequences based on NCBI RefSeq transcripts and human genome build GRCh37/UCSC hg19.

After gene specific filtering, data are analyzed to identify sequence variants and most deletions and duplications involving coding exons; however, technical limitations and inherent sequence properties effectively reduce this resolution for some genes. Alternative sequencing or copy number detection methods are used to analyze or confirm regions with inadequate sequence or copy number data by NGS. Reportable variants include pathogenic variants, likely pathogenic variants and variants of uncertain significance. Likely benign and benign variants, if present, are not routinely reported but are available upon request.

#### **Clinical Sensitivity:**

The clinical sensitivity of sequencing and deletion/duplication analysis of the genes included in the HDCT Panel depends in part on the patient's clinical phenotype and family history. In general, the sensitivity is highest for individuals with clearly defined connective tissue disease and a family history of disease. The technical sensitivity of sequencing is estimated to be >99% at detecting single nucleotide events. It will not reliably detect deletions greater than 20 base pairs, insertions or rearrangements greater than 10 base pairs, or low-level mosaicism. The copy number assessment methods used with this test cannot reliably detect copy number variants of less than 500 base pairs or mosaicism and cannot identify balanced chromosome aberrations. Assessment of exon-level copy number events is dependent on the inherent sequence properties of the targeted regions, including shared homology and exon size. For the B3GALT6 gene, sequencing but no deletion/duplication analysis, is performed. Recombination of TNXB with its pseudogene (gene conversion or TNXB/XA fusion), is not evaluated..

## Test Information Sheet

| Gene              | Protein   | Inheritance                             | Disease Association(s)     |
|-------------------|---|---|----------------------------|
| ACTA2             | ACTIN, ALPHA-2, SMOOTH                                | AD                                      | fTAAD                      |
|                   | MUSCLE, AORTA   |   |                            |
| AEBP1             | AORTIC  | AR                                      | cIEDS                      |
|                   | CARBOXYPEPTIDASELIKE                                  | 7.41.2                                  |                            |
|                   | PROTEIN   |   |                            |
| ADAMTS2           | ADAM METALLOPEPTIDASE                                 | AR                                      | dEDS                       |
|                   | WITH THROMBOSPONDIN TYPE                              | ,                                       | 4200                       |
|                   | 1 MOTIF 2   |   |                            |
| ALDH18A1          | ALDEHYDE DEHYDROGENASE 18                             | AD, AR                                  | Cutis laxa                 |
|                   | FAMILY MEMBER A1                                      | , |                            |
| ATP6V0A2          | ATPASE H+ TRANSPORTING V0                             | AR                                      | Cutis laxa                 |
|                   | SUBUNIT A2  | ,                                       |                            |
| ATP6V1E1          | ATPASE H+ TRANSPORTING V1                             | AR                                      | Cutis Iaxa                 |
| -                 | SUBUNIT E   |   |                            |
| ATP7A             | ATPASE COPPER   | XL                                      | Menkes, OHS                |
|                   | TRANSPORTING ALPHA                                    |   |                            |
| B3GALT6           | BETA-1,3-   | AR                                      | spEDS                      |
|                   | GALACTOSYLTRANSFERASE 6                               |   |                            |
| B4GALT7           | BETA-1,4-   | AR                                      | spEDS                      |
|                   | GALACTOSYLTRANSFERASE 7                               |   |                            |
| B3GAT3            | BETA-1,3-   | AR                                      | Joint laxity/dislocations, |
|                   | GLUCURONYLTRANSFERASE 3                               |   | short stature,             |
|                   |   |   | dysmorphism, CHD           |
| BGN               | PROTEOGLYCAN I  | XL                                      | Meester-Loeys syndrome,    |
|                   |   |   | SEMDX                      |
| CBS               | CYSTATHIONINE   | AR                                      | Homocystinuria             |
|                   | BETASYNTHASE  |   |                            |
| CHST14            | CARBOHYDRATE (DERMATAN 4)                             | AR                                      | mcEDS                      |
|                   | SULFOTRANSFERASE 14                                   |   |                            |
| COL1A1            | COLLAGEN TYPE I ALPHA 1                               | AD                                      | OI, aEDS, rarely cEDS or   |
| 001 / 10          |   |   | vEDS                       |
| COL1A2            | COLLAGEN TYPE I ALPHA 2                               | AD, AR (rare)                           | OI, aEDS, cvEDS            |
| COL2A1            | COLLAGEN TYPE II ALPHA 1                              | AD, AR (rare)                           | Stickler syndrome, SED,    |
|                   |   |   | achondrogenesis, Kniest    |
| COL3A1            | COLLAGEN TYPE III ALPHA 1                             | AD                                      | syndrome<br>vEDS           |
| COLSA I<br>COL4A1 | COLLAGEN TYPE III ALPHA 1<br>COLLAGEN TYPE IV ALPHA 1 | AD                                      | Hereditary angiopathy with |
| COL4AI            |   |   | nephropathy, aneurysms,    |
|                   |   |   | and muscle cramps          |
|                   |   |   | (HANAC)                    |
| COL5A1            | COLLAGEN TYPE V ALPHA 1                               | AD                                      | cEDS                       |
| COL5A2            | COLLAGEN TYPE V ALPHA 2                               | AD                                      | cEDS                       |
| COL9A1            | COLLAGEN TYPE IX ALPHA 1                              | AD, AR                                  | MED, Stickler syndrome     |
| COL9A2            | COLLAGEN TYPE IX ALPHA 2                              | AD, AR                                  | MED, Stickler syndrome     |
| COL9A3            | COLLAGEN TYPE IX ALPHA 3                              | AD, AR                                  | MED, Stickler syndrome     |
| COL11A1           | COLLAGEN TYPE XI ALPHA 1                              | AD, AR                                  | Stickler syndrome,         |
|                   |   | ,                                       | Fibrochondrogenesis        |

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| COL11A2 | COLLAGEN TYPE XI ALPHA 2   | AD, AR | Stickler syndrome,<br>Fibrochondrogenesis   |
|---------|--|--------|---|
| COL12A1 | COLLAGEN TYPE XIIALPHA 1   | AD, AR | mEDS  |
| DSE     | DERMATAN SULFATE<br>EPIMERASE                                    | AR     | mcEDS   |
| EFEMP2  | EGF CONTAINING FIBULIN-LIKE<br>EXTRACELLULAR MATRIX<br>PROTEIN 2 | AR     | Cutis laxa  |
| ELN     | ELASTIN  | AD     | Cutis laxa, Supravalvular<br>Aortic Stenosis  |
| FBLN5   | FIBULIN 5  | AD, AR | Cutis laxa  |
| FBN1    | FIBRILLIN 1  | AD     | Marfan syndrome,<br>Acromicric dysplasia,<br>Geleophysic dysplasia,<br>Weill-Marschani syndrome,<br>Stiff Skin syndrome |
| FBN2    | FIBRILLIN 2  | AD     | Congenital contractural arachnodactyly  |
| FKBP14  | FK506 BINDING PROTEIN 14   | AR     | kEDS  |
| FLNA    | FILAMIN A  | XL     | EDS variant with PVH  |
| LOX     | LYSYL OXIDASE  | AD     | fTAAD   |
| LTBP4   | LATENT TRANSFORMING<br>GROWTH FACTOR BETA<br>BINDING PROTEIN 4   | AR     | Cutis laxa  |
| MAT2A   | METHIONINE<br>ADENOSYLTRANSFERASE II,<br>ALPHA                   | AD     | fTAAD   |
| MED12   | MEDIATOR COMPLEX SUBUNIT   | AD     | Lujan syndrome, Ohdo syndrome, FG syndrome  |
| MFAP5   | MICROFIBRILLAR-ASSOCIATED<br>PROTEIN 5                           | AD     | fTAAD   |
| MYH11   | MYOSIN, HEAVY CHAIN 11,<br>SMOOTH MUSCLE                         | AD     | fTAAD   |
| MYLK    | MYOSIN LIGHT CHAIN KINASE  | AD     | fTAAD   |
| NOTCH1  | NOTCH, DROSOPHILA,<br>HOMOLOG OF, 1                              | AD     | Aortic valve disease  |
| PLOD1   | PROCOLLAGEN-LYSINE,<br>20XOGLUTARATE 5-<br>DIOXYGENASE           | AR     | kEDS  |
| PRDM5   | PR DOMAIN 5  | AR     | BCS   |
| PRKG1   | PROTEIN KINASE, cGMP-<br>DEPENDENT, REGULATORY,<br>TYPE I        | AD     | fTAAD   |
| PYCR1   | PYRROLINE-5-CARBOXYLATE<br>REDUCTASE 1                           | AR     | Cutis laxa  |
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RIN2

| SKI      | V-SKI AVIAN SARCOMA VIRAL<br>ONCOGENE HOMOLOG                              | AD | Shprintzen-Goldberg syndrome |
|----------|--|----|------------------------------|
| SLC2A10  | SOLUTE CARRIER FAMILY 2<br>(FACILITATED GLUCOSE<br>TRANSPORTER), MEMBER 10 | AR | Arterial tortuosity syndrome |
| SLC39A13 | SOLUTE CARRIER FAMILY 39<br>MEMBER 13                                      | AR | spEDS                        |
| SMAD2    | MOTHERS AGAINST<br>DECAPENTAPLEGIC,<br>DROSOPHILIA, HOMOLOG OF, 2          | AD | fTAAD, LDS                   |
| SMAD3    | MOTHERS AGAINST<br>DECAPENTAPLEGIC,<br>DROSOPHILA, HOMOLOG OF, 3           | AD | fTAAD, LDS                   |

Abbreviations: AD – autosomal dominant; aEDS- arthrochalasia Ehlers-Danlos syndrome; AR – autosomal recessive; BCS – Brittle Cornea Syndrome; cEDS-classical Ehlers-Danlos syndrome; CHD – congenital heart defect; clEDS – classical-like EDS; cvEDS - Cardiac-valvular Ehlers-Danlos syndrome; dEDS- dermatosparaxis Ehlers-Danlos syndrome; fTAAD – familial thoracic aortic aneurysm and dissection; JP/HHT – juvenile polyposis/hereditary hemorrhagic telangiectasia; kEDS- kyphoscoliotic Ehlers-Danlos syndrome; LDS – Loeys-Dietz syndrome; MACS

- Macrocephaly, alopecia, cutis laxa, and scoliosis; mcEDS - musculocontractural Ehlers-Danlos syndrome; OHS – Occipital horn syndrome; OI - ostegenesis imperfecta; PVH – periventricular heterotopia; SEMDX – spondyloepimetaphyseal dysplasia, X-linked; spEDS- Spondyldysplasia type Ehlers-Danlos syndrome; vEDS- vascular Ehlers-Danlos syndrome; XL – X-linked

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